

Start crop Point



End crop Point

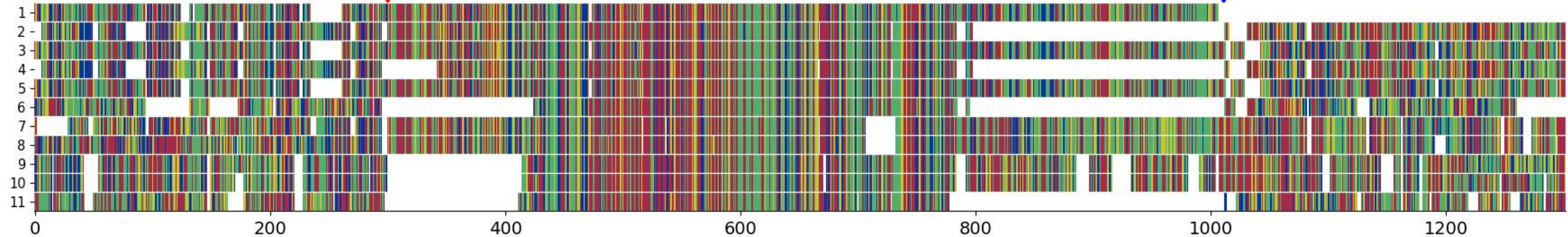


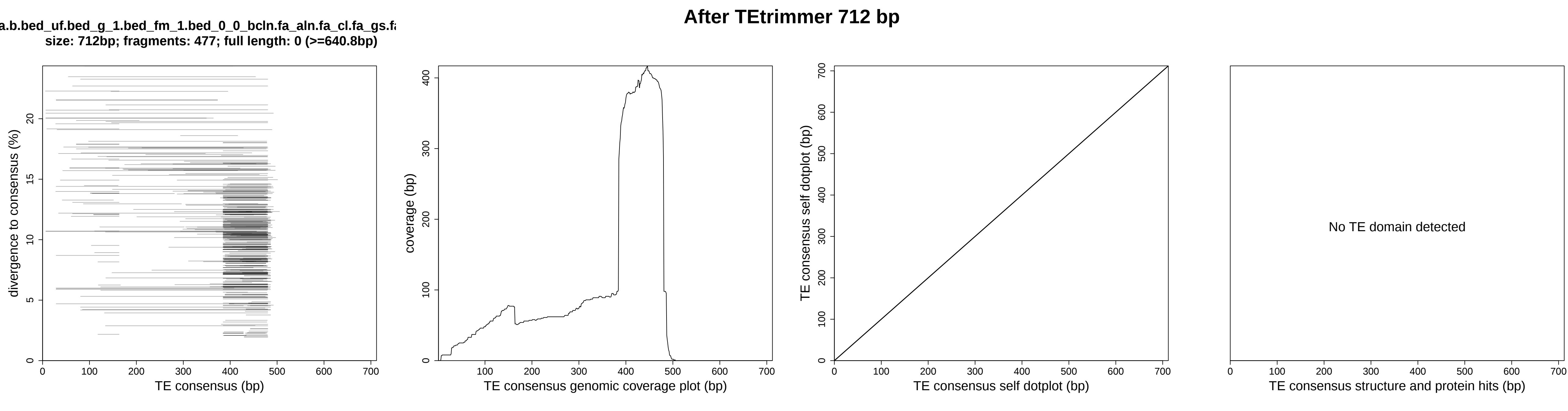
MSA length = 712



Start crop Point

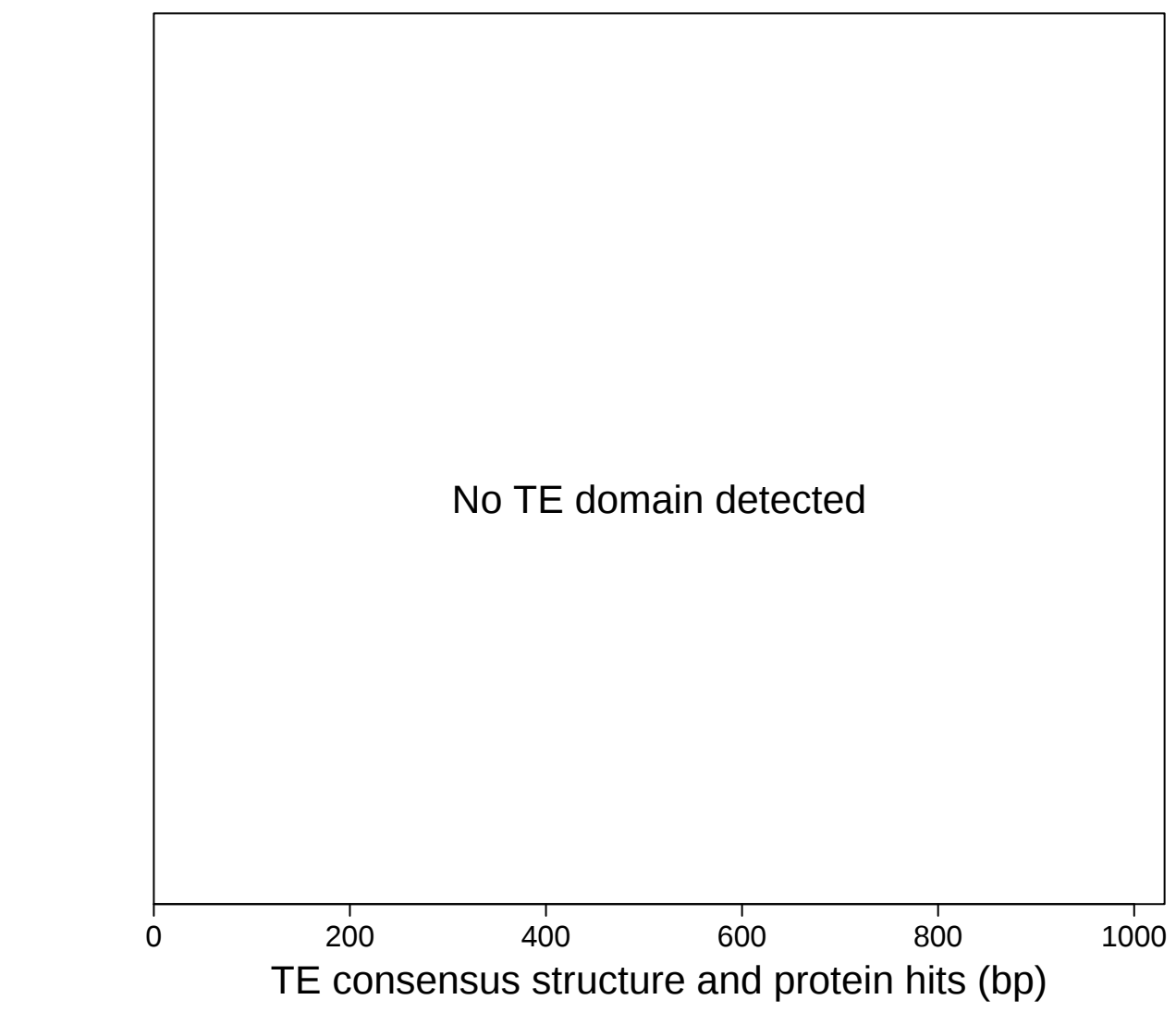
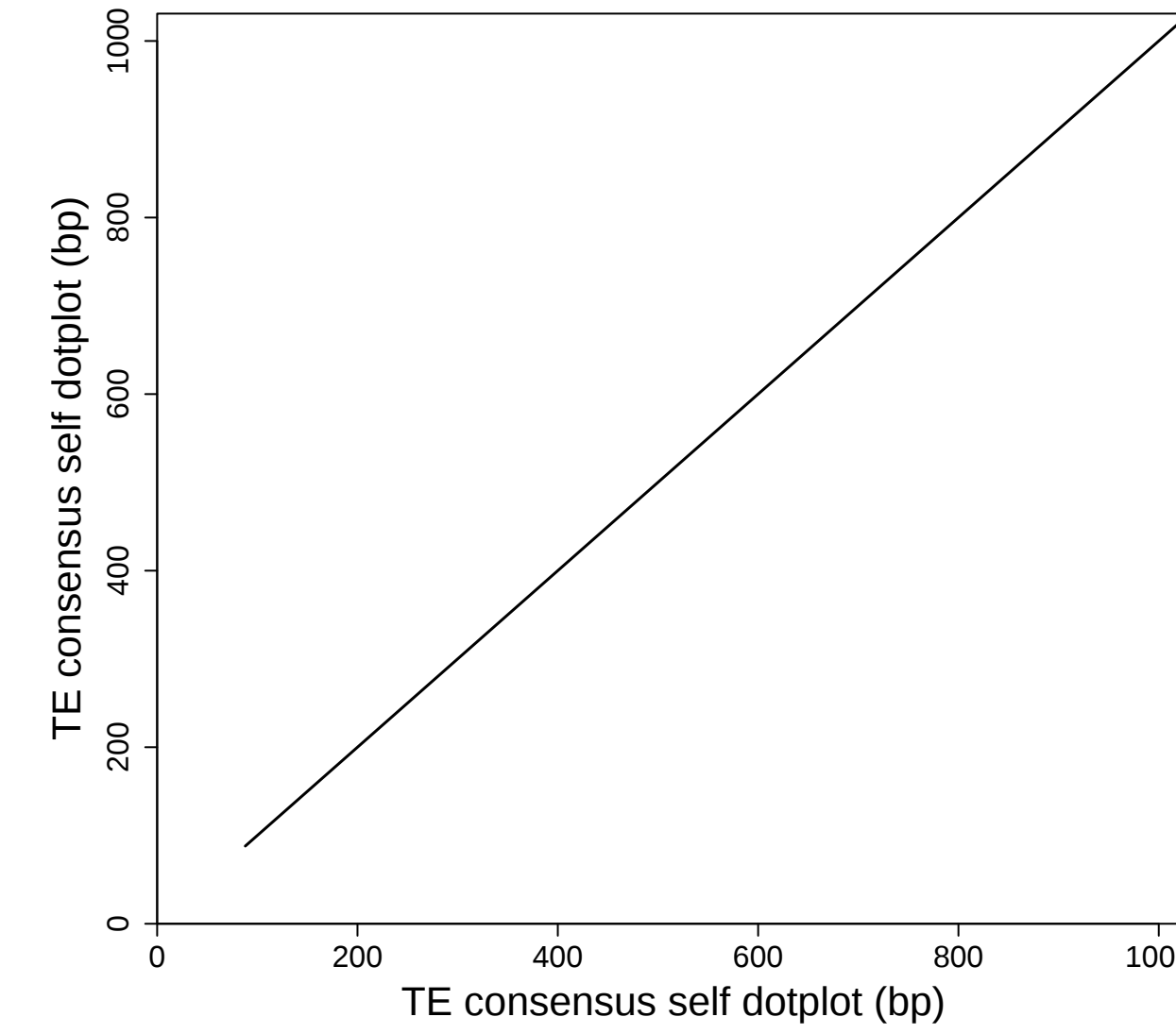
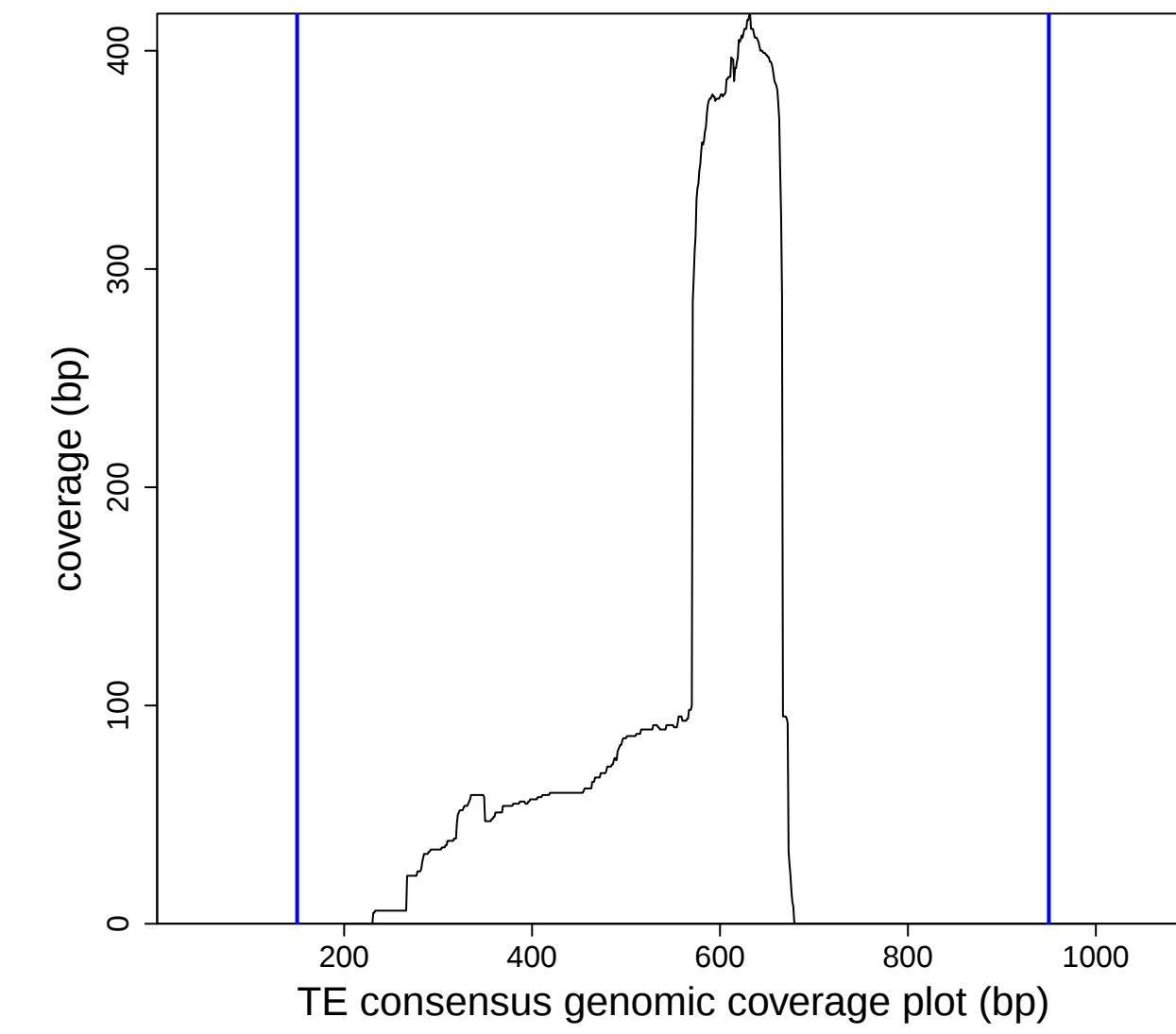
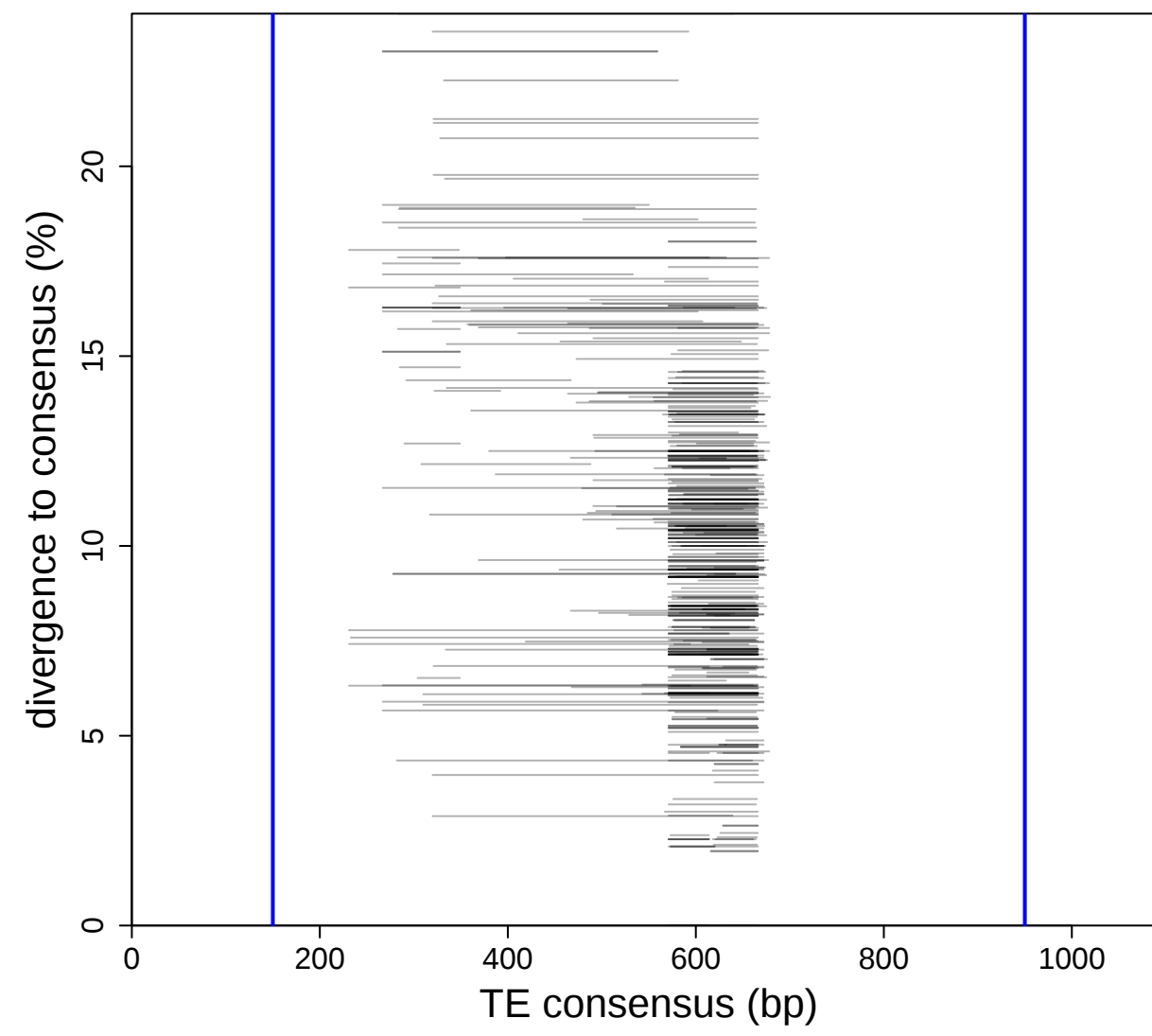
End crop Point



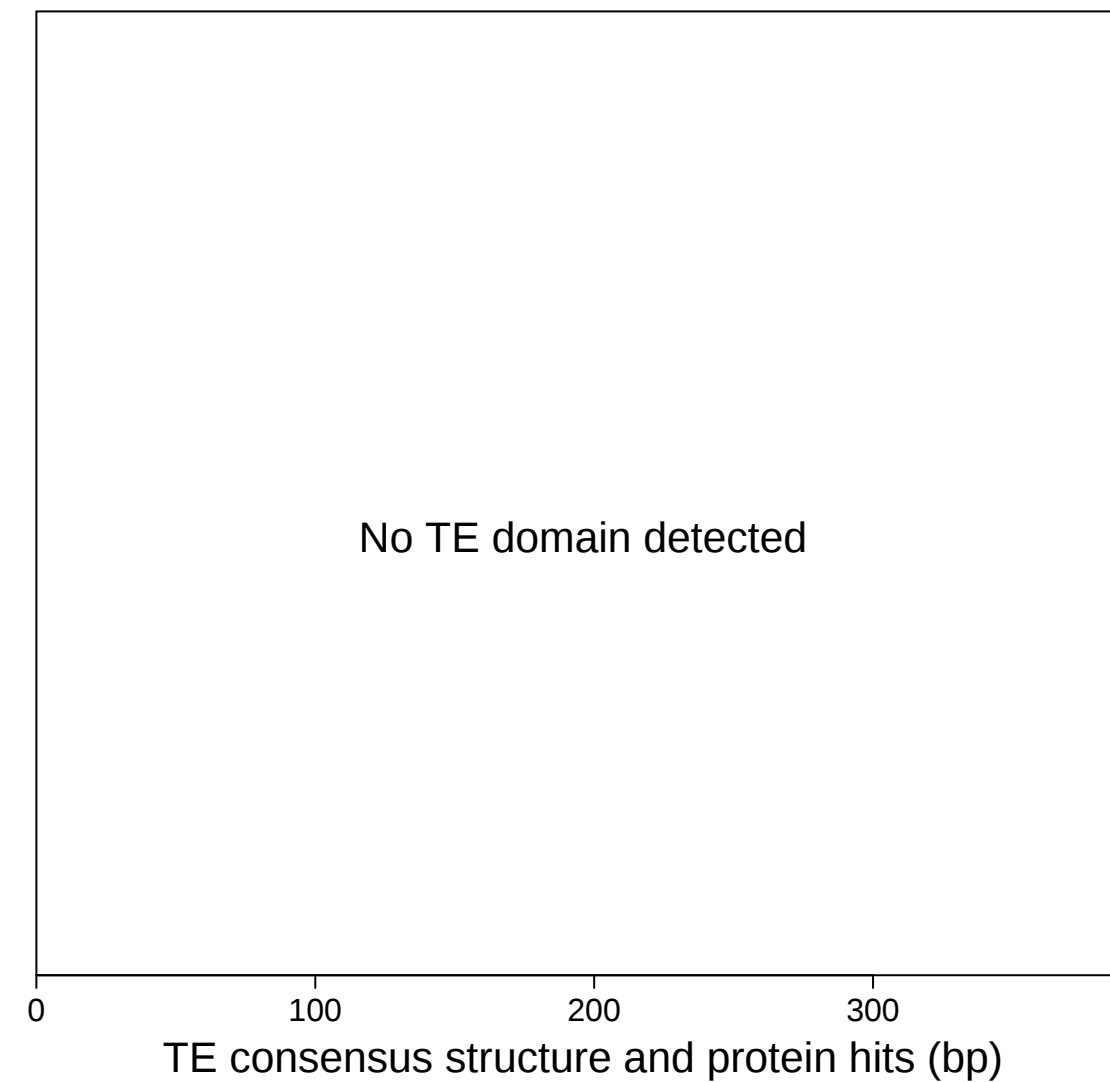
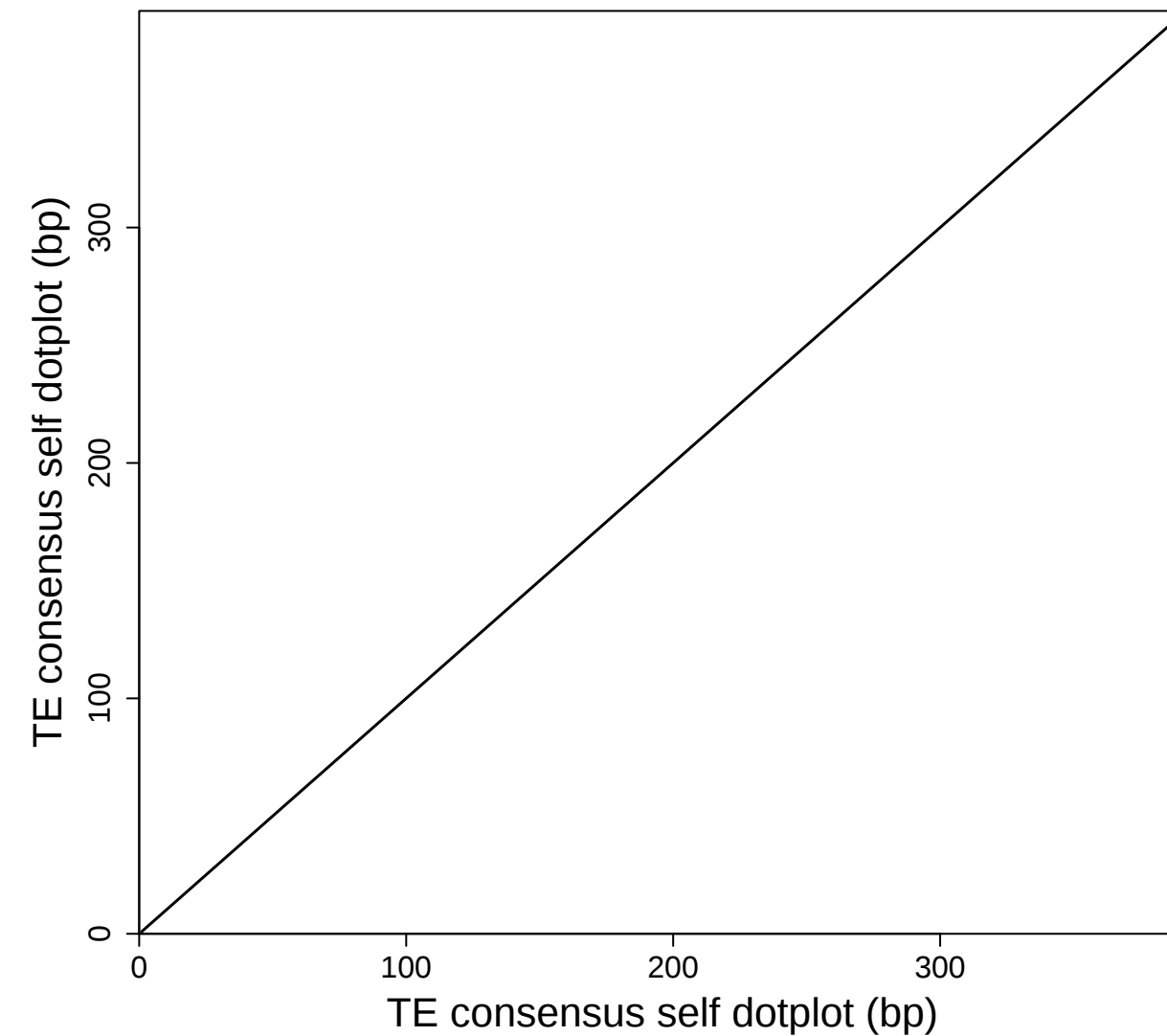
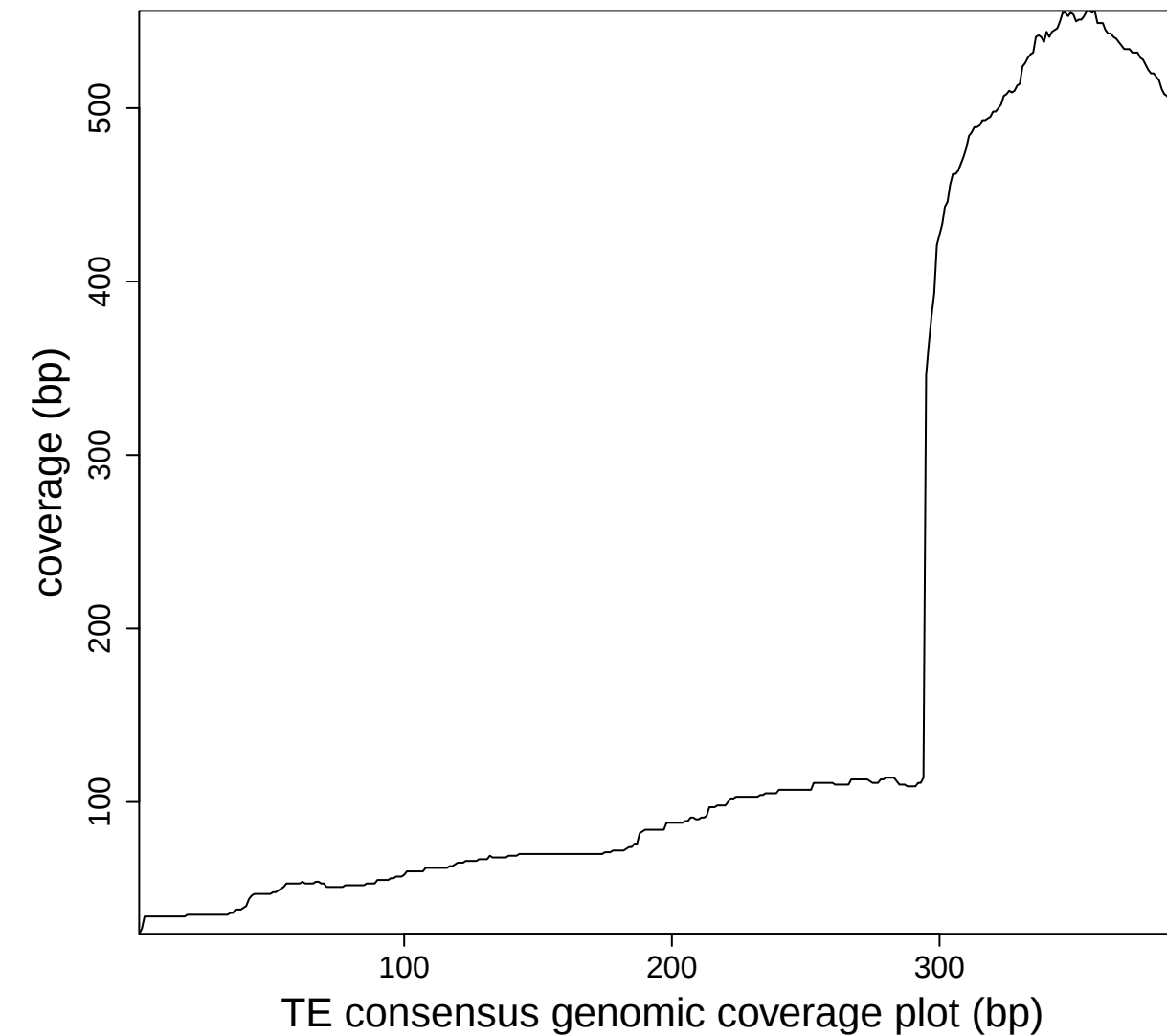
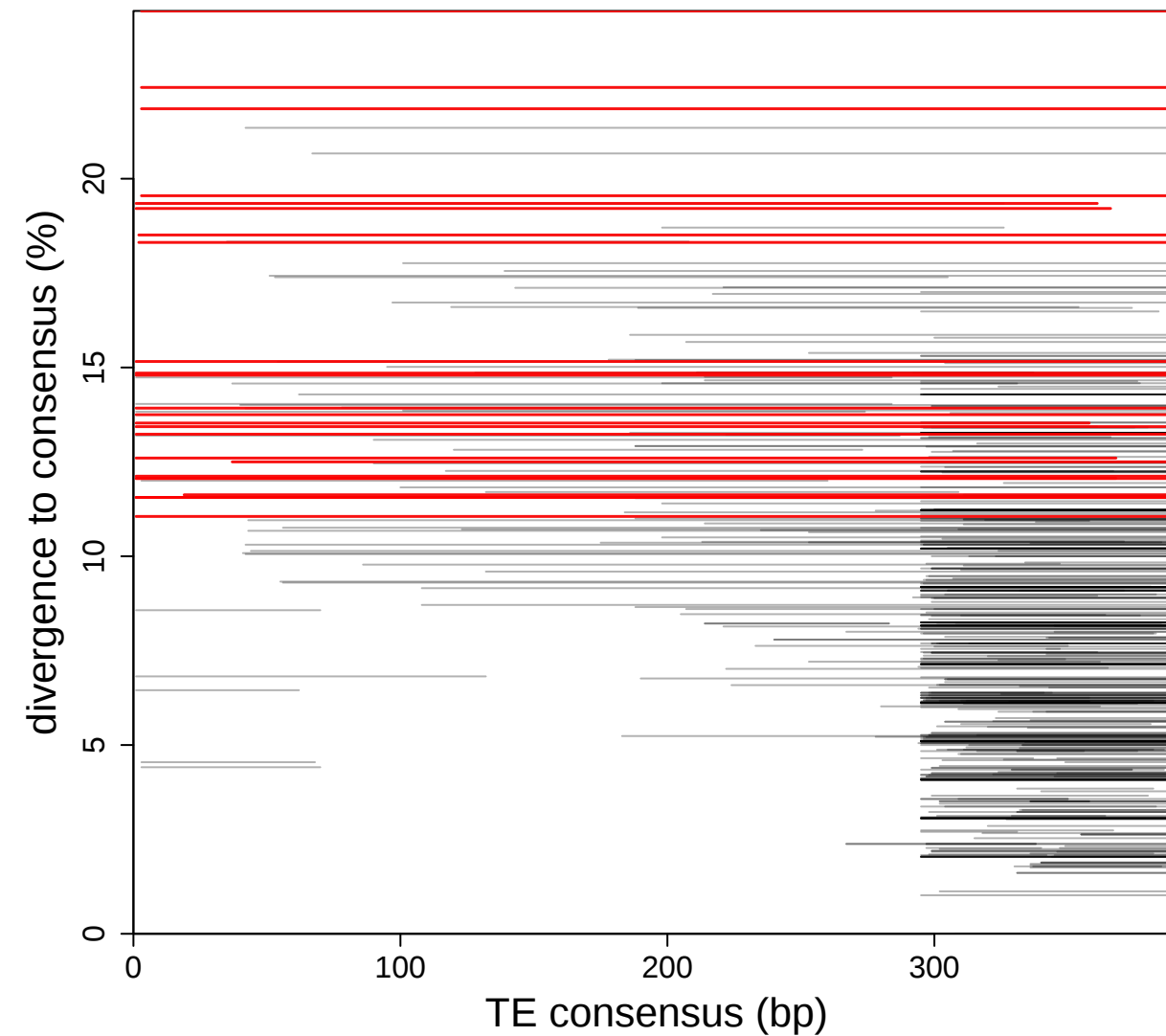


2.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 1100bp; fragments: 461; full length: 0 (>=990bp)

After TEtrimmer Extended plot Blue lines are boundaries

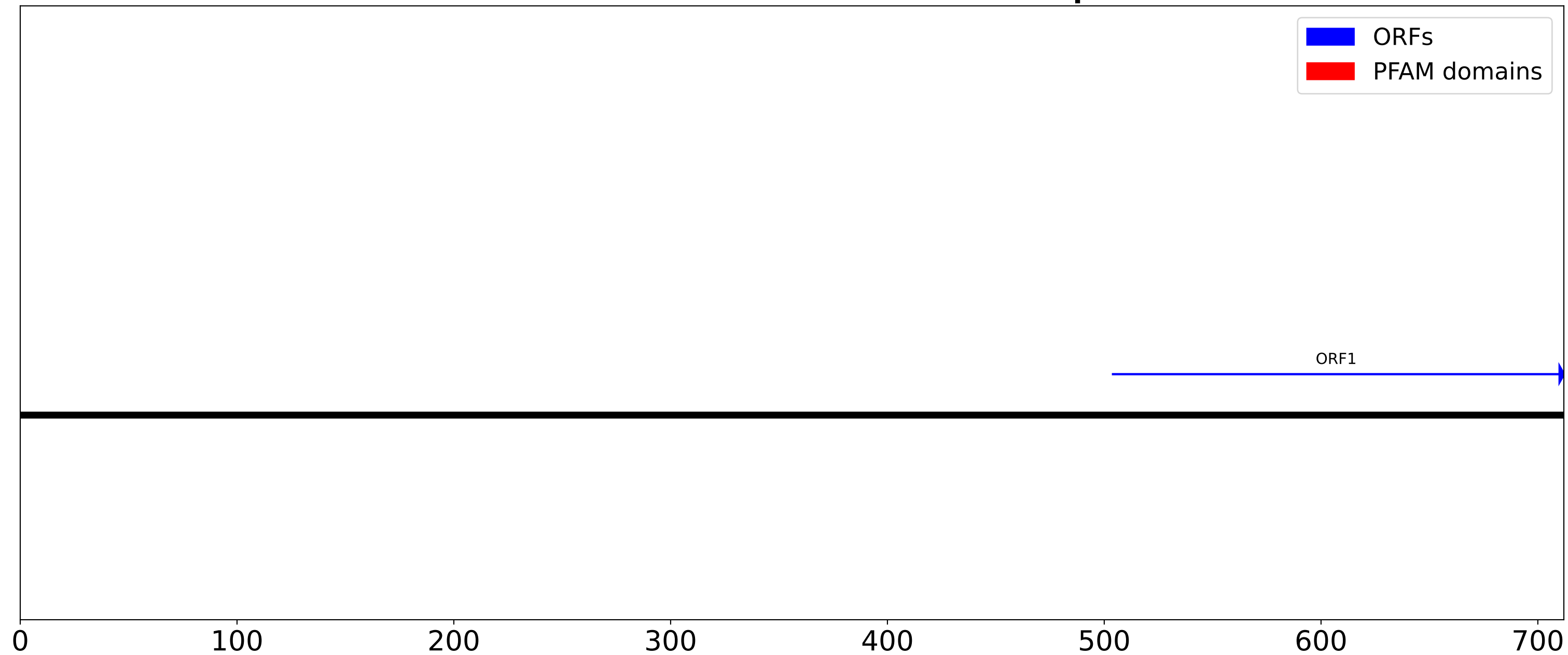


TE: rnd_5_family_1182
size: 392bp; fragments: 599; full length: 25 (≥ 352.8 bp)

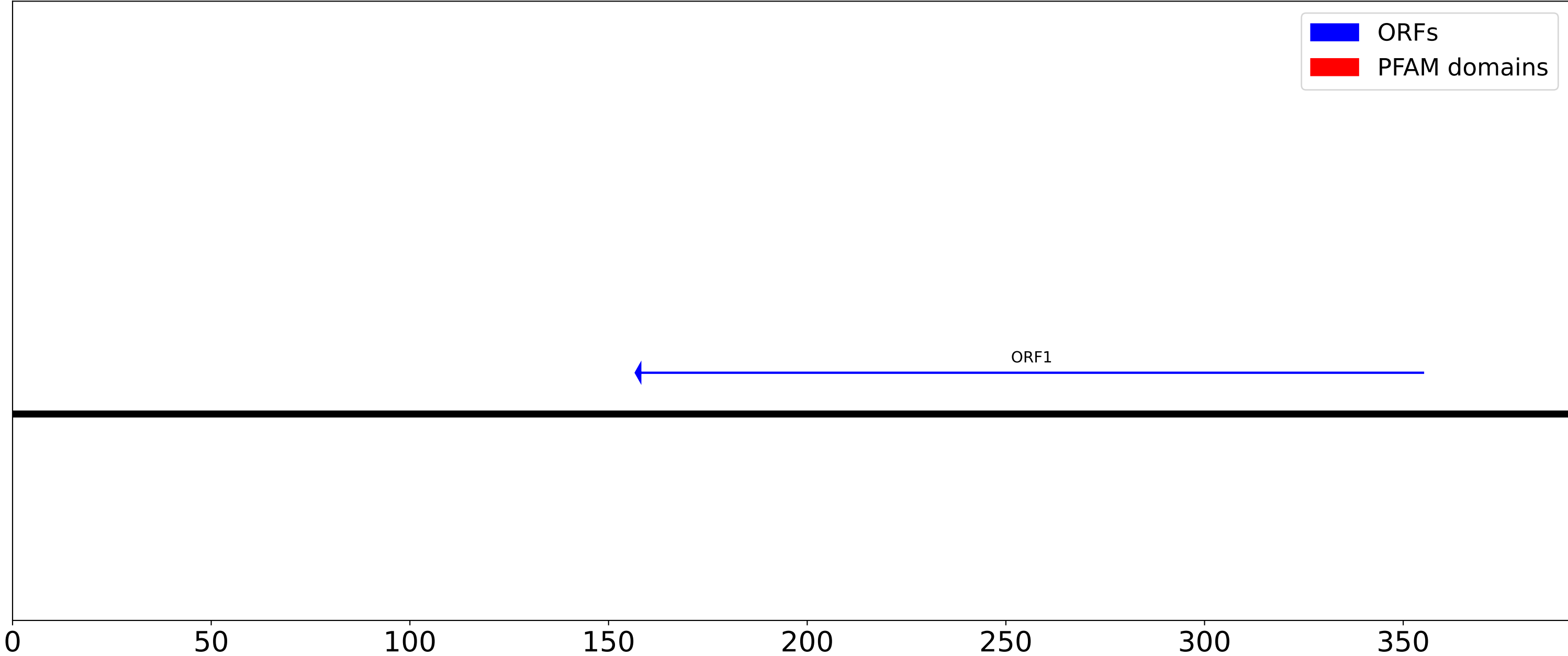


Before TEtrimmer 392 bp

After TEtrimmer ORF and PFAM domain plot



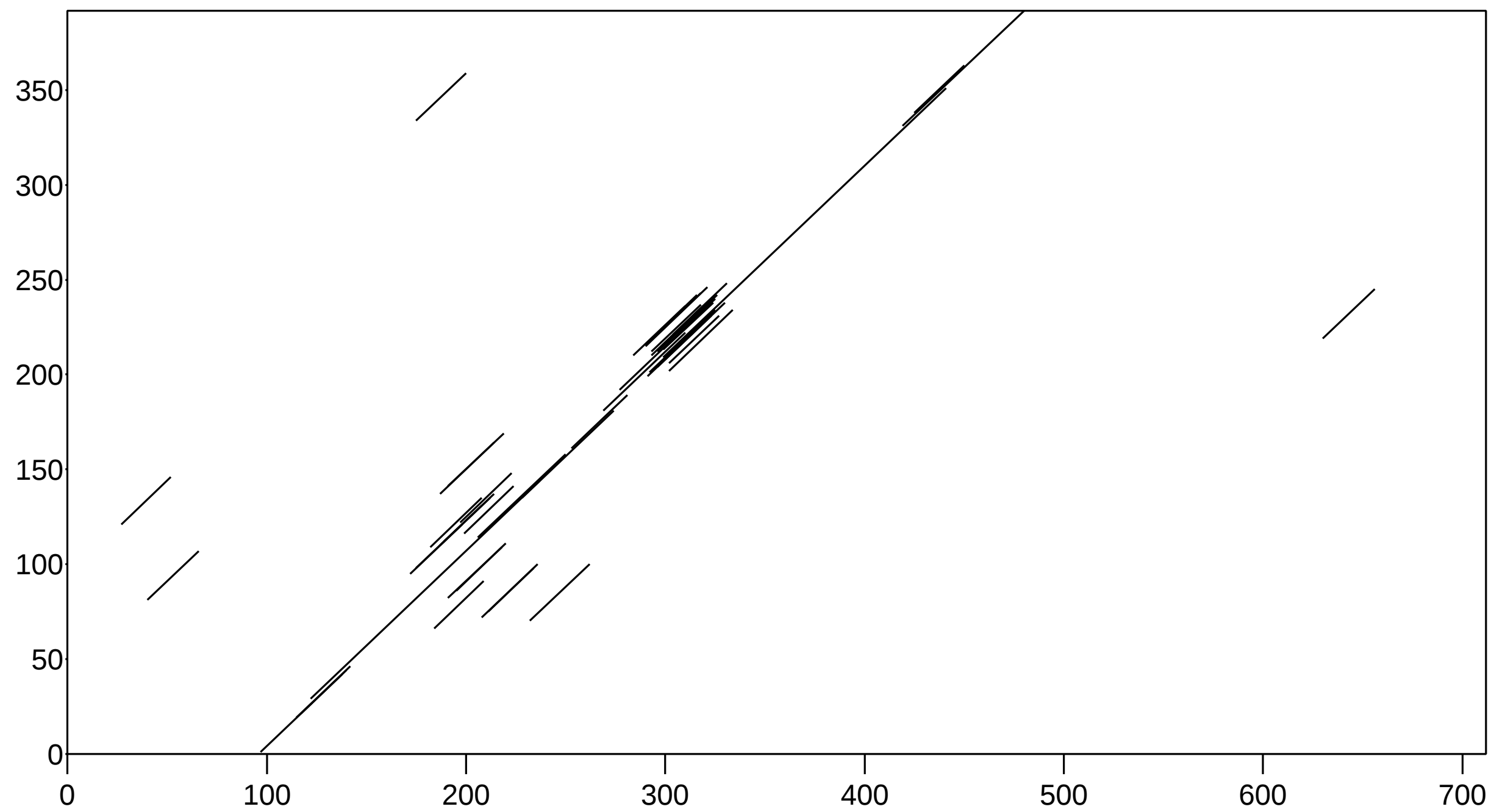
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 10/10/25)

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)